### IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of:GAN et al

Art Unit:

Serial No.To Be Assigned

Examiner:

Filed: March 14, 2001

Atty. Docket: CL0001165

For: ISOLATED HUMAN RAS-LIKE PROTEINS, NUCLEIC ACID MOLECULES ENCODING THESE HUMAN RAS-LIKE PROTEINS, AND USES THEREOF

# SUBMISSION OF SEQUENCE LISTING UNDER 37 C.F.R. § 1.821(a)

Honorable Commissioner of Patents and Trademarks Washington, D.C. 20231

Sir:

In compliance with 37 C.F.R. § 1.821(a), applicants submit the Sequence Listing, including the paper copy of the Sequence Listing and the computer readable copy of the Sequence Listing.

### In the Specification:

The Sequence Listing is provided on pages 56-72 of the specification in the above-identified application.



## **REMARKS**

In accordance with 37 C.F.R. § 1.821(f), the paper copy of the Sequence Listing and the computer readable copy of the Sequence Listing submitted herewith in the above application are the same.

It is respectfully believed that this application complies with the Sequence Listing requirements and is now in condition for processing.

Respectfully submitted,

**CELERA GENOMICS** 

3y: \

Justin D. Karjalá

Date: \_March 14, 2001\_\_\_\_\_

Celera Genomics Corporation 45 West Gude Drive, C2-4#20 Rockville, MD 20850

Tel: 240-453-3067 Fax: 240-453-3084 SEOUENCE LISTING

```
<110> GAN, Weiniu
<120> ISOLATED HUMAN RAS-LIKE PROTEINS,
  NUCLEIC ACID MOLECULES ENCODING THESE HUMAN RAS-LIKE
  PROTEINS, AND USES THEREOF
<130> CL001165
<160> 8
<170> FastSEO for Windows Version 4.0
<210> 1
<211> 2349
<212> DNA
<213> Human
<400> 1
tecteegqte gecegeete ggggeageta gtggegeage ceceegeeeg eggeeetgge 60
ctcccqqqcq qcqcqqcaqq qqaqqqqtta aqctqccqca qqqaccqccq cqtqcqqqqc 120
gagagggage ceeeggtggg ggtggegeag ceggeggggt teggteegag eeeggtggga 180
ggetecegga gegeageetg ggeeeageee acceegegee ggeggeeatg geaggeacee 240
tggacctgga caagggetgc acggtggagg agetgeteeg egggtgeate gaageetteg 300
atgacteegg gaaggtgegg gaccegeage tggtgegeat attecteatg atgeaecect 360
ggtacatccc ctcctctag ctggcggca agctgctcca catctaccaa caatcccgga 420
aggacaacte caatteeetg caggtgaaaa egtgeeacet ggteaggtae tggateteeg 480
cetteceage ggagtttgae ttgaaceegg agttggetga geagateaag gagetgaagg 540
ctctgctaga ccaagaaggg aaccgacggc acagcagcct aatcgacata gacagcgtcc 600
ctacctacaa gtggaagcgg caggtgactc agcggaaccc tgtgggacag aaaaagcgca 660
agatgteeet gttgtttgae caectggage ceatggaget ggeggageat eteacetaet 720
tggagtateg etecttetge aagateetgt tteaggacta teacagttte gtgacteatg 780
getgeactgt ggacaacece gteetggage ggtteatete ectetteaae agegteteae 840
agtgggtgca gctcatgatc ctcagcaaac ccacagcccc gcagcgggcc ctggtcatca 900
cacactttgt ccacgtggcg gagaagctgc tacagctgca gaacttcaac acgctgatgg 960
cagtggtcgg gggcctgagc cacagctcca tctcccgcct caaggagacc cacagccacg 1020
ttagccctga gaccatcaag ctctgggagg gtctcacgga actagtgacg gcgacaggca 1080
actatggcaa ctaccggcgt cggctggcag cctgtgtggg cttccgcttc ccgatcctgg 1140
gtgtgcacct caaggacctg gtggccctgc agctggcact gcctgactgg ctggacccag 1200
cccqqacccq gctcaacqqq gccaagatga agcagctctt tagcatcctg gaggagctgg 1260
ccatggtgac cagcctgcgg ccaccagtac aggccaaccc cgacctgctg agcctgctca 1320
eggtgtetet ggateagtat cagaeggagg atgagetgta ceagetgtee etgeageggg 1380
agcogogoto caagtootog coaaccagoo coacgagttg caccocacca cocoggococ 1440
cggtactgga ggagtggacc tcggctgcca aacccaagct ggatcaggcc ctcgtggtgg 1500
agcacatcga gaagatggtg gagtctgtgt tccggaactt tgacgtcgat ggggatggcc 1560
acateteaca ggaagaatte cagateatee gtgggaactt ceettacete agegeetttg 1620
gggacctcga ccagaaccag gatggctgca tcagcaggga ggagatggtt tcctatttcc 1680
tgcgctccag ctctgtgttg ggggggcgca tgggcttcgt acacaacttc caggagagca 1740
actecttgeg eccegtegee tgeegeeact geaaageeet gateetggge atetacaage 1800
agggeeteaa atgeegagee tgtggagtga actgeeacaa geagtgeaag gategeetgt 1860
cagttgagtg teggegeagg geceagagtg tgageetgga ggggtetgea eceteaceet 1920
cacccatgca cagccaccat caccgcgcct tcagcttctc tctgccccgc cctggcaggc 1980
gaggetecag geetecagea atececetee cageagagat eegtgaggag gaggtacaga 2040
cggtggagga tggggtgttt gacatccact tgtaatagat gctgtggttg gatcaaggac 2100
tcattcctgc cttggagaaa atacttcaac cagagcaggg agcctggggg tgtcggggca 2160
```

ggaggctggg gatggggtg ggatatgagg gtggcatgca gctgagggca gggccagggc 2220

<210> 2 <211> 615 <212> PRT <213> Human <400> 2

<400> 2
Met Ala Gly Thr Leu Asp Leu Asp Lys Gly Cys Thr Val Glu Glu Leu
1
Leu Arg Gly Cys Ile Glu Ala Phe Asp Asp Ser Gly Lys Val Arg Asp
20
Pro Gln Leu Val Arg Ile Phe Leu Met Met His Pro Trp Tyr Ile Pro
35
Ser Ser Gln Leu Ala Ala Lys Leu Leu His Ile Tyr Gln Gln Ser Arg
50
Lys Asp Asn Ser Asn Ser Leu Gln Val Lys Thr Cys His Leu Val Arg
65

70 75 80

Tyr Trp Ile Ser Ala Phe Pro Ala Glu Phe Asp Leu Asn Pro Glu Leu
85 90 95

Ala Glu Gln Ile Lys Glu Leu Lys Ala Leu Leu Asp Gln Glu Gly Asn 100 105 110

Arg Arg His Ser Ser Leu Ile Asp Ile Asp Ser Val Pro Thr Tyr Lys
115 120 125

Trp Lys Arg Gln Val Thr Gln Arg Asn Pro Val Gly Gln Lys Lys Arg 130 135 140

Lys Met Ser Leu Leu Phe Asp His Leu Glu Pro Met Glu Leu Ala Glu 145 150 155 160

His Leu Thr Tyr Leu Glu Tyr Arg Ser Phe Cys Lys Ile Leu Phe Gln
165 170 175

Asp Tyr His Ser Phe Val Thr His Gly Cys Thr Val Asp Asn Pro Val 180 185

Leu Glu Arg Phe Ile Ser Leu Phe Asn Ser Val Ser Gln Trp Val Gln
195 200 205

Leu Met Ile Leu Ser Lys Pro Thr Ala Pro Gln Arg Ala Leu Val Ile
210 215 220

Thr His Phe Val His Val Ala Glu Lys Leu Gln Leu Gln Asn Phe

225 230 235 240
Asn Thr Leu Met Ala Val Val Gly Gly Leu Ser His Ser Ser Ile Ser 245 250 255

Arg Leu Lys Glu Thr His Ser His Val Ser Pro Glu Thr Ile Lys Leu 260 265 270

Trp Glu Gly Leu Thr Glu Leu Val Thr Ala Thr Gly Asn Tyr Gly Asn 275 280 285

Tyr Arg Arg Arg Leu Ala Ala Cys Val Gly Phe Arg Phe Pro Ile Leu 290 295 300

Gly Val His Leu Lys Asp Leu Val Ala Leu Gln Leu Ala Leu Pro Asp 305 310 315 320

Trp Leu Asp Pro Ala Arg Thr Arg Leu Asn Gly Ala Lys Met Lys Gln 325 330 335

Leu Phe Ser Ile Leu Glu Glu Leu Ala Met Val Thr Ser Leu Arg Pro 340 345 350

Pro Val Gln Ala Asn Pro Asp Leu Leu Ser Leu Leu Thr Val Ser Leu 355 360 365
Asp Gln Tyr Gln Thr Glu Asp Glu Leu Tyr Gln Leu Ser Leu Gln Arg

```
370
                        375
                                            380
Glu Pro Arg Ser Lys Ser Ser Pro Thr Ser Pro Thr Ser Cys Thr Pro
                    390
                                        395
Pro Pro Arg Pro Pro Val Leu Glu Glu Trp Thr Ser Ala Ala Lys Pro
                405
                                    410
Lys Leu Asp Gln Ala Leu Val Val Glu His Ile Glu Lys Met Val Glu
            420
                                425
                                                    430
Ser Val Phe Arg Asn Phe Asp Val Asp Gly Asp Gly His Ile Ser Gln
                            440
Glu Glu Phe Gln Ile Ile Arg Gly Asn Phe Pro Tyr Leu Ser Ala Phe
                        455
                                            460
Gly Asp Leu Asp Gln Asn Gln Asp Gly Cys Ile Ser Arg Glu Glu Met
                    470
                                        475
Val Ser Tyr Phe Leu Arg Ser Ser Ser Val Leu Gly Gly Arg Met Gly
                485
                                    490
                                                        495
Phe Val His Asn Phe Gln Glu Ser Asn Ser Leu Arq Pro Val Ala Cys
            500
                                505
                                                    510
Arg His Cys Lys Ala Leu Ile Leu Gly Ile Tyr Lys Gln Gly Leu Lys
                            520
Cys Arg Ala Cys Gly Val Asn Cys His Lys Gln Cys Lys Asp Arg Leu
                        535
                                            540
Ser Val Glu Cys Arg Arg Arg Ala Gln Ser Val Ser Leu Glu Gly Ser
                    550
                                        555
Ala Pro Ser Pro Ser Pro Met His Ser His His Arg Ala Phe Ser
                565
                                    570
Phe Ser Leu Pro Arg Pro Gly Arg Arg Gly Ser Arg Pro Pro Ala Ile
                                585
Pro Leu Pro Ala Glu Ile Arg Glu Glu Glu Val Gln Thr Val Glu Asp
                            600
Gly Val Phe Asp Ile His Leu
    610
<210> 3
<211> 20951
<212> DNA
<213> Human
<400> 3
acagaaaggt cctgtttcta agtcttacat taccaagact gaggtgcggg ggcggtcctg 60
gatececege eccaaggetg ggagggeae geeteggaag ggaggtttgg ggteggtggt 120
ttcacagtga gtgtgtctga agccaaatgg tcggaaaccg ttacccgctc tcctaggccc 180
ggctagtggg gaccccaacc gcctgcggct gcccctccca agttcctccc tqttqqccag 240
gcatccaggt ctccagtctc cgagctqcqq aqaacccacc gccacatgcq gctgccctt 300
tecattegae cetgtgggga gecaggette eggggeeceg tteeteetgt gtgaaetggg 360
cccccgccc ccattcccag acatcaaggc cgcgtctcca gatagccacg atttcattcc 420
tegetececa caggiecete tecceaaaat atteceatet tgteetagee cateececag 480
actateteaa ggaccagetg tecceaegee eeegacetee actaggeetg tgecaeeege 540
tgeetgeagg aagaegeeeg gteeegggee gggttageee catgggaaeg gtttgteteg 600
aaaacaggaa cccgagctgg gggctgggcg gggcgccct tccccaccgc agtccgcttc 660
etgeceetee eggetteete egeeegacae eeaggeaggg eggggggcae tggggegtee 720
geggttgggg gaggggetet tegttteggt eeceeteee gegteeeggg eggegggee 780
teeggtegee egeetegggg cagetagtgg egeageeece egeeegegge cetggeetee 840
egggeggege ggeaggggag gggttaaget geegeaggga eegeegegtg eggggegaga 900
gggagecece ggtgggggtg gegeageegg eggtgeggag eteegegeag gggeggaggg 960
gggaggggc agcetggege gggggeggg geggggegge ggggageggg geegeggegt 1020
```

ggagageggg egggageege ageeggageg aggeeggegg gegggagege aeggaggtgg 1080

ggtcggccag gccggtgcgg gctccttgcg gcaggtccca agagtgagtg ggcgagcgcg 1140 ggcggggcgc caggcgaagg agggcgcgc ccccagcgac tcccccccg cccagggcgg 1200 cgcgggcggg ctggggggg cgagcgggtg gggagtctgc ggcccgggtc tgggagaggg 1260 ggcagcggcc acgagagcta aggcgcgctg gatccccgga gggcggagga cctccacggt 1320 geacceaget tttcccagec acettecage ggggecetee eeegegtace eccatttggc 1380 agatgagaaa attgaggctc ccagaggcca agtgattctc aaggtcacac gaggaagcgg 1440 tagaqccaqq cqqqqacqqc tctqqqtqqc tcttagqaaa agtccqcctg agaactccqt 1500 acaggagete ceetgteete cageetgggg gagtgagtat gtgtagggee ggggtacett 1560 tccqtqqqqc aaqqctctqc caaaatctqq qaqtqaqqqq aqtcagqqaq ctqqqqccqc 1620 agggcgggcc ctgcaccqca aatgggaggg gggcqacqga atgggcgtgc gcacccatgg 1680 qqqtqtqtqc atqtqtqtqq qaqtqtacat qcqtqqaqaq gcactqcctt qcqttqtqc 1740 acacgtgtga ggatgtcagc gcctgtgtgg ccgcgggact caaggctggc ctggctcaag 1800 tgaacagcac gtccaggagg cgacctcgtc cgcgggtttg cattctgggg tggacgagct 1860 gggtatgtgt gcctgagggt ttcttcgtgc aggtgtgcac agggtgtggg tgccattgtg 1920 tgtgagagac ggaggatggg gaggccggtg cctgtggccc ggtgcgtgta agtgcggacg 1980 cctgcacctc cacttaggtc cccggcctcc gacgactaac ttgggtgtgg agtgtttgcc 2040 cctgccaggg tgcgtatgac cccgccagtg accggagttg ctaatggtgt catgcaccca 2100 coggecacce ttggcgcgag cgccccctc tggacaccct gctccgtgcg cgctcacagt 2160 tegeetgtge ggggeegggg ceagggteag gageegggga tagggaggaa gagggeetgt 2220 ggacaagetg ageegggace eetgggacet ttgeggaggt ggeetgggag egeteagtte 2280 ccaggetgag getteceget gaegeeteet ggeegeageg ggeteeeee geeeeaggaa 2340 tgttcctctc ccatccagtc cgcctcccct agggcaggcc ccctgggggc tgccgcagcc 2400 cegectegee tteetggget eeegggaggg ggegaggega geaggaegee tgggttetet 2460 coccccacct cocataccag ggagaaattc ctccgaggtc ccctcaggct ctgggttccc 2520 aaaataaccc tgcgggggaa gggaggctgt ggagggaggg aagcgggagg ggcgcagagc 2580 cgagctgcgg gqtqctqcaq gtgcctctqg ggagagggcg cgaggagaag gcgccctgcg 2640 gggggctggg cgccagccag tcctgggatc ttggttcgtc cccatcctcg tgaagcccct 2700 eggeetteee gegaeteega gggtgggeeg gaageetete tgegggteeg ttteeeaact 2760 ggcgggttgc accatcccgg gccagaccgt ttaaccccgg gagtggccgc gggggacaac 2820 tecgececty tecageaggg ggegtgeegg eccegececg tttetgeeeg eggggeeget 2880 ceccegeceg egacteegea gacteeeget etgeetetee egggacaggg gtteggteeg 2940 agcceggtgg gaggeteceg gagegeagee tgggeecage ceaeceegeg eeggeggeea 3000 tggcaggcac cctggacctg gacaagggct gcacggtgga ggagctgctc cgcgggtgca 3060 tegaageett eggtgagtgg etegggaggg cacaeggage etgageetag eeeegagtet 3120 gagccegggt cectgeetee caggcacagt ceagggcaca gecetgaece ggacceacee 3180 tgctccgcag cgtgcagtct ctttaacgaa agcctcctcc gcaacgcagg gcagagagat 3240 geacgecett cagacagatg aggttteect tetetageet teeccagegg eggegaaggg 3300 agggccgggt cccggactct gacacttgag gggcattatc tgtctcccgg ggaatccgga 3360 ggaactcgct atctccggcc tgggagctgt ttccggctaa tggggggcgg cttatctggt 3420 gaaggggtgc cccttccccc caagcgctca ggaaatgacc tctggattct tgaccccqqg 3480 gaacccagge teetteegee eeagetggtt eeeeteegga egatgggegg etegggeget 3540 cccctcctcc agtcctcagg gcgtgcctat ctctcgccca ccacaccttt cctctctaat 3600 ttgcctcctg ctctcggagt cctgggcaag caggaggtgg gcggggtcga gcgtgcaccc 3660 gaaggaccga tacctggcgg gttgcggggt gaggatgagg catggtagct gcggacccag 3720 ctcagccacc tgtctttgac ccttcggagt cagatgactc cgggaaggtg cgggacccgc 3780 agctggtgcg catgttcctc atgatgcacc cctggtacat cccctcctct cagctgqcgq 3840 ccaagctgct ccacatatat ccttcgccgg ccttgccaag gcccccgccg tcggagccca 3900 tgcgcagccc ctctgcccag cccaggtgca gaatgagcct cgctcctaag tataggccac 3960 tecttatece agageteagg egtegteeca geeteeaact agggeetagg etetgeece 4020 tecttgetce tagegacteg gteetgteec caggetetgt ceceaqeeqa ggeettgeec 4080 tecttetece tagagtetag ggeetgeece tgetteagge ttgggtgege eeegtgeate 4140 tetetetece agageceagg etttgettte ageeteette ageacetagt eeteeacece 4200 cacctccaac ccctcccaga gctcaagcct cacccccagc atctccgcag agcgcaagcc 4260 ccatccctag aacgtgtctc ctagaaccag gcccgcccc cagcctccct ccacgcaggc 4320 ctccctttct agagttaagc ggcctcctta accetctcct tcacctacca acaatcccgg 4380 aaggacaact ccaattccct gcaggtgaaa acgtgccacc tggtcaggtg agtctttccc 4440 ctggggctct agcccctccc ctttctccct tctctctggc ttcaggctgg cctggaggag 4500

qqqqcaqqqc qctqtttctq qqaqtqqqtt tgaaccctqq cttqtccqqq tqqqcaqtqc 4560 tgccacagge teacceette etgggtetgg geettaattt tettttetge geagtgeggg 4620 tggttgtctc aagggtctaa tgtacacttg gagtggcgaa ggaaagagct ggaaccatag 4680 tttgagggtc tttttgctta ggtgactata atctcaaata gctccttgca gcctgctggg 4740 tgatqqtqqq qqaaqqqcta tcttqqqtqa ctccccqctc ctccaqqtac tggatctccq 4800 cetteccage ggagtttgae ttgaaccegg agttggetga geagateaag gagetgaagg 4860 ctctgctaga ccaagaaggg aaccgacggc acagcagcct aatcgacata gacagcgtgt 4920 gcgtgggggg agcacagagg gctggggggg cactcagtat cctataccat ctgtgcttaa 4980 taaatgtctg ttgaactgaa tgagtgaggg tcatgttgct ctctcgctta aaaaccttcc 5040 atggctccct attgccttca acatgcctcc tctgggcage ttggcgttcc tgcctcatct 5100 tocactgoca coaccoatco cacacaceto etectgtage tgegetgggt eggeteeceg 5160 teggetgage tetegagtee titeteatea tggtgetetg eteatateat ecceettget 5220 gcctcctccq tqttaccaaq actcaqttca qqcatqaaqt ctccqtqqqc tctqaqqqtt 5280 eggggetett eeggggtaga atttgtegtt eccaectetg titteeatgg eactitigtae 5340 agactectgt acaaagacet etgtacatgt gteacgetgt tttgtgatea tgtgtttetg 5400 tgtctgtctc cctcagtaga ctgtgagctc ctcgagggca ggaaccgtgt cttactcatc 5460 tetgtattee cagegeetag cacagtgeet ggeacagagt aegttgttea taaatgtgtg 5520 ttgagtgcat gacggggtgg ggggagatga ggaggagttg ctgggactgg gaacattcgt 5580 gcctaggaca gtgcctcgca ttatgtaggt tctcagtaag cgtgaatggt gtgtctgtgt 5640 gagtgggggg ccacgaggca tgcgcatgtc cagcaaaggg ctcactaccc ctgcccccc 5700 agccctacct acaagtggaa gcggcaggtg actcagcgga accctgtggg acagaaaaag 5760 cgcaagatgt ccctgttgtt tgaccacctg gagcccatgg agctggcgga gcatctcacc 5820 tacttggagt ategeteett etgeaagate etggtgegge eegagggetg gggggteagg 5880 ggtccaatgt gggctggaag agagttctag gaggggcagg gtccctggcg taggctgggt 5940 cacagggtgc atcaggggtt tcagtgtaac cactgaaggt cagctggagg gtgaggagtg 6000 gctatcagtg aggggagagg ccggcaaggt gctgaggcca ctcctcatgc ccccagtttc 6060 aggactatca cagtttegtg acteatgget geactgtgga caacceeqte etggageggt 6120 teateteect etteaacage gteteacagt gggtgeaget catgateete ageaaaceea 6180 cageceegea gegggeeetg gteateaeae aetttgteea egtggeggag gtgeetgeee 6240 ctccctcccg gtgtctccca accacccac atgccagtca ggccaaccct tcccttcccc 6300 taacccactg cottototot agataagotg ggccaaatto tgggcccact cagtgactoc 6360 ctgcctctcc gtccccattt gccttccaga agctgctaca gctgcagaac ttcaacacgc 6420 tgatggcagt ggtcgggggc ctgagccaca gctccatctc ccgcctcaag gagacccaca 6480 gccacgttag ccctgagacc atcaaggtgc ctgggactgg ggaggggccg gtgcttccca 6540 ggtctgtctt cactgggtcc tcccagcagc actgggggct gggcacagct gtcctcattt 6600 gatagatatg gaaatggagg ctcagagggg ttaagtgctt ttctcagttt gcacaatggc 6660 aacagcagag tgggggctca caggtcgtca gggaccccaa agctagtact ttttttttt 6720 tttttaagac agggtetete tetetgttgt eeagactgga gtteagtggt geagteacaa 6780 gctcactgca gccttgaatt cctgagctca atcgatcctc ccacctcagc ctcctgagta 6840 gctgggacta caggtgtacg ccaccatgcc taatttttgt attgttatta atttttttt 6900 ttttttttta gagatggggt tttgccatgt tgcccagact ggtcttgaac tcctgggctc 6960 aagtgateeg eetgeettgg eeteecaaag tgetgagatt atggettgag eeattgtgee 7020 ttgccacttg tagtttcttc ttttctttct ccttcatttt ttattatttt tgaagtattt 7080 tgaagtattg agtaacatac atatagaaaa gtatataaaa acatatgaga ctgggcgtag 7140 tageteacae etgtaateee ageaetttgg gaggetgagg tgggeagate aegtgacate 7200 aggagtttga gaccagcctg gccaacaagg tggaaaccca tctctactaa aatacaaaaa 7260 ttagccaggc atggtggcac gcacctggaa tccaagctac ttgggaggct gaggcaggag 7320 gagaattact tgaactcagg aggcggaggt tgcagtgagc caagattgtg ccacttcact 7380 ccagcctggg cgacagagtg agactccatc taaaaaaaaa gaaaagtata taaaaacata 7440 tgaatagttt aaaqaaaaat tgtaaaqaaa acactgtgta actactgccc gggttgggaa 7500 atagaacett gecaggeece caagegeeca geaetttaga geataactee eteeceaega 7560 cttttgcaat gatgatettg cttttettta tagetteace atgtaggtat geggtecaaa 7620 tattttatgt cttgcttttt tcattccaca tggttctgag agtcttttca ttctgtcatg 7740 tggagcaatt gtttttcat tttcattgcc atataatatt ttattgtacg tctaccccaa 7800 ttcatttatt tatttatttt tttgagatgg agtctgtctc tgtcatccag gctggagtgc 7860 ggtggcgaga tctcatcact gcaacttccg tctcctgggt ttacgtgatt ctcgtgcctc 7920

agectectga gtagetggga ttatgggete gtaceaecae gtetggetaa ttttttgtag 7980 agacaggett teaccatgtt geogaggetg gtettgaact cetgagetea ggeaateeac 8040 ccqctttaqc ctcccaaaqt gctqqqatta caqqtqtqaq ccactqcccc cagcctaccc 8100 caatttatgt attgattcta ttgttgaatg ttggggtttt tccttttctt ttctttcttt 8160 ctttttcttt cttttttct tttttttgga gagggagtct tgctctgtcg ccaggctgga 8220 gtgcagtgac gctaatttgg ctcactgcat cactgcaccc tctgcctccc gggttcaagc 8280 gattetectg ceteageete etgagtaget gggaetaeag geatgeacea ceaeaceegg 8340 ctaatttttg tatttttta gtagagatga ggtttccacc atgttggcca agatggtctc 8400 catctcttga cctcatgatc catctgccat ggcctcccaa agtgctgaga ttacaagtgt 8460 gagccaccac gcccagctgg tttttccagt ttttgctgtt tggacggggt ggctgagtat 8520 gttcttccag gtcattgtcc tgtgctgcct tgcctccctg agcctctgtt tctcctgtta 8580 aatgttgatg attccctgca tccaggcctg gtttagaggt gtggtgcttt tggcagtgag 8640 tattgccttg aattcatggc aatgaattca atccccaggg gctgagagag ccagtcgtgg 8700 gggacagtaa gggaggtttt tactctttca cctqtccctg accctgactc ctcctcaccc 8760 cctcctacat ttccagggct gaggtaggga ggatagttgt gggggtatga ctcctctgtc 8820 ctttgtcccc agctctggga gggtctcacg gaactagtga cggcgacagg caactatggc 8880 aactaccggc gtcggctggc agcctgtgtg ggcttccgct tcccgatcct gggtgtgcac 8940 ctcaaqqacc tqqtqqcct qcaqctqqca ctqcctqact qqctqqaccc aqcccqqacc 9000 cqqctcaacq qqqccaaqat qaaqcaqctc tttaqcatcc tqqaqqaqct qqccatqqtq 9060 accagectge ggccaccagt acaggccaac eccqacetge tqaqectget caeggtqagg 9120 agcagggggc agggaggtgg ggagctgggc accaggggtt gacagtttcc ccaggtcctg 9180 gctgtgggcg tggcctgggg ctctgggttc tggccaagaa actgagatct agcgtgggct 9240 ctggggtttg gagtggatgc tgagaagggg tccaggctct ggttggggct gtggactgag 9300 gtetgatete caggetggta tgtggaetgt gggeagtttg aactgggeet gggteeeggg 9360 ttgagttctg gcaatgggct gtgttctagg gctgggccaa gctctgcatt ctgtgggcag 9420 gggtggtttc taagcatggc cctgggctcg gagtgaagtt ctgggcttgg ctttacactt 9480 ggtcttgggg tctagggtgg gagttgggtt ctggtttaga tccagacaag gttctagaca 9540 ttqqqctqqq qcttaaqtqt taaqqtttqq aqtqqattct taqctqcttc tqqqctctqq 9600 aggggatcag ggttgaaatc agagettetg getgggttee gaeetggett etteeetgae 9660 atcttggcaa tatgttgtgt tcaaggtttg gggccatgct gtggtttgat ctgtgcgctg 9720 ggatgacatq ggggttgctg tgctgtgttc taagccaggc tttgtcctga gtctagcttc 9780 tgaccogage tetggetgag etgtggeete taggtegace tittggeeetg ggetetgtgg 9840 ccgtgggcag gggccagtgg gggtgatcag atctgtgtgt cccaggtgtc tctggatcag 9900 tatcagacgg aggatgagct gtaccagctg tecetgeage gggageegeg etecaagtee 9960 teggtgaggg ggtacteect cetetecaet etgecettee etectgagaa teccaggatg 10020 tgaggatggg aagagctett agcagccace teacceatee atettgtagg acagaggeat 10080 cctgggggta gggcagtagt gttgggcaga cttccctctc ccagggattc ccctctctqt 10140 teccegggge tetgggetee ecetgeetet ggeeetaget eaggeeegae eattteeata 10200 gecaaccage eccaegagtt geaccecace acceeggeec eeggtactgg aggagtggae 10260 ctcggctgcc aaacccaagc tggatcaggc cctcgtggtg gagcacatcg agaagatggt 10320 ggaggtgagc tcctgcggag cctgagcagt gtgtggggag aggccagttt gccggagcac 10380 tgccctggaa gccagcacga gtgtcctgtt caagacccag cactcagccc ctaggagtca 10440 cagggcctgg caggccagct gcacggggct gaagtgcccc tgggtagggt gggggtggag 10500 gtatggaacg ggggtggtgt cagagacctc tctgagacac acctcatcaa atggactggg 10560 aacgtgggaa gggacaggac ctgatgtccc ctttactctc ccctcttctg gctctgcgtg 10620 tecetetgeg tgeeceagte tgtgtteegg aactttgaeg tegatgggga tggeeacate 10680 tcacaggaag aattccagat catccgtggg aacttccctt acctcagcgc ctttggggac 10740 ctcgaccaga accagtgagg agggctgggg acctggggga gagggaaggc aactcagccc 10800 acttctgcct gggcttcagt ttcttgtgtg caagatgagg tcactgagcc agatgatctt 10860 ggcctgggaa gctgccagtg tgggaaaggg cacttgcttt tgtggggagg agaggctgcc 10920 agctgtggag gcgcagtggt atctcacaaa ttcagacaga tggggggctc cacctgagtc 10980 ttgcaaagac tgtgacctgg ggactgtggc tacaaaagtg ctgttttatt tgtggagctc 11040 acagetgtea agaagtgtgg geaacttgag eteetggata gtetgtteta atgaatagat 11100 aagaaaggtt tgtaattagc agtacccagt tgtttatcaa cagttcatat gctgacaatt 11160 tggaaaaaca gctggttctc tgaagtaggt taaacatgcc ccctgaagcc agattcatgc 11220 cctatttttg ctgagcagaa aaaactccat tcaaaattta aagtccatct caggtcgatt 11280 tattttttaa tgttacctgt atttcaaaaa tctgttgttt tttatttcca cattacaaaa 11340

atccacggta aaataaaatc taggtggtaa aataaattta tagtgaacaa aatgtttaaa 11400 gtaagaagtg agaggccagg tgcggtgcct cacgcctgta atcctagcac tttgggagac 11460 tgagttggca ggatcaattc aggccaggag tttgagccca gcctgggcaa cagagtaaga 11520 ccctqtctct acaaaaatta ttattattat ttttgagaca gagtctcact ctgttgccca 11580 qqctqqaqtq caqtqqtaca atctcqqctc qctqcaacct ccacttcctq ggttcaagtg 11640 attetectge tteagettee tgagtagetg ggattaeagg catgeateae egtgeetgge 11700 taatttttgt atttttagca gagatggggt tttaccatgt tggccaggct ggtctcaaac 11760 tettgacete aagtgateta eetgeettgg eeececaaag tgetaggatt acaggeatga 11820 gctactgctc ctagcctaaa aaaatttttt ttgggcatgg gtggcacgtg cctgtagtcc 11880 cagetactea ggaggetgag geaggaggaa eeettgagee eaggaggttg agaetgeagt 11940 qaqctqtcat cacaccactg cacttcagcc tgggtgactg cgcgagatca cccccatcaa 12000 aaaaaaaaaa aaaagaaaaa aaaaggaaga aatgaaagtc ccctctttcc ttttccactg 12060 gtagaagttg ccatgattaa gcactgttaa caatattaag cttggcagta tgtggattct 12120 tecagtette tttteecagg caggtgeaca ttgatagaga ttttgtttgt ttggtgtetg 12180 tttcatggac aaacaggatt agagcataaa tctagttctg cttgtggctt ttatcatagc 12240 tgctttattt cttctcccag attttaggca gaggtagttg agttccatgt tttctccctg 12300 qqttqqtqqq tqqattttta tctaqaccac cttttcaqtq agaatgaccc tttqaqacqa 12360 tggaggeete agetteatge agegggetea geettaacce tecaceteet geaggeecea 12420 ggaaagcccc tggttgggta tcaaaaacct agcacctggt tcggcaggag ggagaccagc 12540 accggctccc caggaccagg cccagctcac cacttcattg taaagctccc tctttgtttc 12600 tqqaacttqq qtqtttccat ttctttctta caaaattatc tatqcattta caqcaattqt 12660 tgatatatet ttaggeagea tetaggtaet tgtagtgggt tetetttttt etttttett 12720 ttttttaatc accetetett ttttttgaga eagagtetea etetgteget eaggetggag 12780 tgcaatagcq cgatcttggc tcactgcaac ctctgcctcc caggttcaag taattctcat 12840 gcctcagcct cccaaqtaqc tgaqattaca ggcactggcc accagacccg gctaattttt 12900 ttttcttttt ctttttttg agacggagtt tcgctctttg ttgcccaggc tggagtacag 12960 tggtgtgatc tcggctcact gcaacctccg cctcccgggt tcaagtgatt ctcctgtctc 13020 agceteecga gtagetggga ttacaggege gegecaccat geetggetaa ttttgtattt 13080 tttttttttt gagacagagt ctcactctgt cacccagact ggagtgcggt ggcgcgatct 13140 cggctcactg caagctctgc ttcccgggtt catgccattc tcctgcctca gcctccggag 13200 tagctgggac tacaagcacc caccaccgtg cccggctaat tttttgtatt tttagtagag 13260 acggggtttc accgtggtct cgacctccag acctcgtgat ccactagcct cagcctccca 13320 aagtgctggg attacaggcg tgagccacct cacccagcct aattttgtat ttttagtaga 13380 gatggggttt caccatgttg cgcaggctgg tattgaactt ctgacctcag gtgatccgcc 13440 cgcctcqqcc tcccqaaqtt ctqqqattat aggcgtgaqc caccgcacct ggcctaattt 13500 ttgtattttt agtagagatg gagttttacc ttgttggcca ggctggtctt gaactcctga 13560 cctcacctca ggtgatctqc ccacctcgqc ctcccaaagt gctgggatta caggcatgag 13620 ccactgtgca cccggcctaa aaatcaccat cttgacagaa cttcacgcct tgctttttgt 13680 tttttttcat ctttgtgctt gttttccact taaccettga tcacagacat ctttccatgt 13740 ggattcatgt agaactacct cattcgttag aacagctgca gagtattcca ctgtgcggtt 13800 agtecateat tteectaace atcetectge tqatqqacag ttaqactqtt ccagttttte 13860 agtatgattc tatgccaggc tgccatgaac qtccttttac tgatccactc aggccagtat 13920 ttctqtaqqa qaaattccta qaaqtqqqat aattqqatca aaaqatatqc acattctaaa 13980 ttaggagaga gactgccaaa ctgacctcag acaaggttgt accagtttgc acccccatca 14040 gcagcgtaca agtgcctgct tcccaacttc ctcgccaaca gggatgctat aaaaagcttc 14100 acaattttgc cagtctcatt ggcaaatggt atcttggtta aatttgcatt tctttaatac 14160 taagtggggg tagggtatet tttcatatgt ttattggcca tttatttctt ctgtcaattg 14220 cctgttctga ttccttgtcc attattctac tgggtttgtt ggtctttttc tcattgattt 14280 ttagaatctc tgttaatgga tattaaccct ttgctgttga atgtgtttgc aaatattttc 14340 tccctgtctg tcatttatgt gtctttttcc atataaattt aaaaaatttt ggtgggctca 14400 ataggtcagt ctttcccttc cgggcttctg ggatttgtgt tcggggtaga aaggccctca 14460 gtgtcttgcc atgtcaccca ggctggagtg cagtggcatg atcttggctc gctgcaacct 14580 ccacctccca ggttcaagtg attetcgtgc cttagcctcc cgagtagctg ggattatagg 14640 tgcctgccac tatgcctggc taattttttg tatttttagt agagacgggg ctttgccatg 14700 ttggccagge tggtettgaa etectgaeet egtgateeae eegeettgge eteceaaagt 14760 gctgggacta caggcgtaag ccactgtgct cggccctata tttttttcag atagccagtt 14820 atectaatge teeettgatt tqatqqacca eetqqatcac acattatqaq eeeectcata 14880 agcaggtggg agtctcaagc gagggccagt cccgatggga atagcacttg gtggctgagg 14940 accetectat etgtgeagae aetgttgtaa aactteacat geateateta atttagteet 15000 caccaaaatc ctatgaaatg taggaatgat cattacaccc atttatagat aaggaaacgg 15060 agggacaggg agattactcc gctacaggtc aagaggcagg gaagtagagc tgcgatttga 15120 actgaggtct gtgtctagaa cacgtgctca ttctttccct aaaatgtatt cataggtgaa 15180 aaagggette tgeggaaage cetgggttat gtgggaaace etggatttae agetgtettt 15240 ccagcaggat gatgcaggag agagaggat qcgatttctc ccaatctctc ctggtcccag 15300 aactcattaq aqaqttctcc ctgctgaqqq ctcccqactq qtqttqcaca cagtacactt 15360 cgggagcccg aggctgatgg ttccatggaa agtacacagt cattttagtt tgcacaccaa 15420 qtqtgaaqtq qqcaqqacaq qccactqttc tqaqaaqqaa cccaqqqaaa qqqactqqcc 15480 caagaccaca cactggttag cggcacttcc cacatctgcc tgacccctag tccagtgccg 15540 ccttttcttt actctgcaac aggagtccaa aatcaggagt tccatgagga cactgggaac 15600 agtgggatgg gttaggccag cggtggatgg ttctggggag ggcccgagct gaagcgcccc 15660 cgcaactccc cacagggatg gctgcatcag cagggaggag atggtttcct atttcctgcg 15720 ctccagctct qtgttqqqqq qqcqcatqqq cttcqtacac aacttccaqq aqaqcaactc 15780 cttgcqcccc gtcqcctqcc qccactqcaa aqccctqqtq aqaqtccctt tcccqqctca 15840 eggeecaage caegeecete eageecegge eeegeeetee ettetggeee egeetetgee 15900 agagecette teaagecagg aaaacetggt aattetattt geeteteete etgtggttet 15960 gcccggggcc ctgaggcggg ctctaaagcc ctagtctcac cctcaagaag gaagaagtag 16020 agteateace tetaaateee teeteecace acggeecete etetattgea gateetggge 16080 atctacaagc agggcctcaa atgccgaggt gagatggaat gactggaagg gctgctgggc 16140 agtgtttttt ttgtttgttt gtttgtttgg gagagttact attttggtgg ggcaattgcc 16200 aaggagtgaa gtaccttaaa atcagaggcg catggccggg catggtggct caagcctgta 16260 atcccagcac tttgggaggc cgaggcgcgc agatcacctg aggtcaggag ttcaagacca 16320 gcctgaccaa catagcgcaa ccccgcctct actaaaaata caaaaagtag ctgggcgtgg 16380 tggcacccac ctgtaatccc agctacttgg gaggctgagg catgagaatc gcttgaacct 16440 gggaggcggg gtttgcagtg agccgagatc acgccactgc actccagcct gggcaacaga 16500 gagggetetg teteaaaaaa aaaaaacaac aaaaaaacce ecaaaaccaa aaccecacaa 16560 aatcagaggc tcaagatgac tgatgtgaag ggagtggcgt ttaagaggcc atttattttg 16620 atgacgcagc tgcccaggaa cagagaacat gggagaaggc atagactgac aattaggagg 16680 aggagaacac tttggaagga gactettatt ttggtgggge agetgeteag gaacaaaggt 16740 tcctggtagg ggggcgcaag cctgcgggat gggatggagg gtattctgac caatgtccct 16800 ggetggetet ceatttgete teeeccagee tgtggagtga actgeeacaa geagtgeaag 16860 gategeetgt eagttgagtg teggegeagg geeeagagtg tgageetgga ggggtetgea 16920 costcaccet cacccatgca cagocaccat caccgogoet toagettete totgeccoge 16980 cctggcaggc gaggctccag gcctccaggt aagagggagt cattctgtac tggcctgtgg 17040 agggaaggat gcagggctac tggggcaaag aacgcaggat ggaagccatt ccaaagtgca 17100 taattotott tttgtggtgg gataataaag aagggacagg ccgggcgcgg tggctcacgc 17160 ctgtaatccc agcactttgg gaggccgagg cgggcggatc acgaggtcag gagatcgaga 17220 ccatcctggc taacacggtg aaaccccatc tttactaaaa atacaaaaaa aaaaaattag 17280 ccaggcqtqq tqqcqqacqc ctqtaqtccc aqctacttqq qaqqctqaqq cagqaqaatq 17340 gcatgaaccc gggaggcggg gcttqcaqtq aqccqaqatc qcqccactqc actccaqcct 17400 gggcgataga gcaagactcc gtctcagaaa aaaaaaaaat aaaaaataaa gaagggacag 17460 gtaagggtgc cagaaagtgg ccaggaagcc ctggaccttc tgaggctgag gagagagacc 17520 ctaatttata aagaggtata aaagtgaaag aggcttcaag attccagtta cagtcttatt 17580 ttgttggagg ggttaacaaa ggattggaga aggtgttata tgagccattg gcttgccttt 17640 ccctttctgg ctgctctgga ggctcttctg qggaaagtcc cttqccctga taatgtcctg 17700 gcagctctct tggggtattt gatggtttta ggtcagtttg ctgaatgaca actggccaaa 17760 tgattatttt gctgagaaca gtccgaacaa ctatgttaaa ctggggtcta aggtagttga 17820 tcacaactgt ttgggttggc ataagtcctc aaaaaacaga ggcaggcaca gggcatacat 17880 cctcaaaaat agaaaagata aatccatttg cattgagcct tccagaagtg ctggggtcta 17940 aaatgtgaaa tacacacaaa attgacattt aagcaaactg cgctgacaaa tctgtqqctq 18000 aaaaaqctgt ggcaaaacaa aaacatagaa aaagagcctc aaaaattggg ctgaggccag 18060 gcatggtggc tcacgcctgt aatcctagca ctttgggaag ccaaggtggg tggatcaccc 18120 

```
aaatacaaaa attagctggg cgtggtggca ggcgcctgta atcccagcta cttgggaggc 18240
tgaggcacga gaatcqcttg aacctgggag gtggaggttg cagagagccg agattgcgcc 18300
aaattgggct gtgaggtcat gcagggaatt gatttttggt gggtgggtct gcttctggga 18420
tgatqtggat gcctcccgtg gagaggggaa gggttgatga agtcccaggg acctggaagt 18480
gtgttctgca gcaatccccc tcccagcaga gatccgtgag gaggaggtac agacggtgga 18540
qqatqqqqtq tttqacatcc acttgtaata gatqgtgagt cctcccacag ctggcaccag 18600
agetececae tgagggetgg gggggagetg gggagtatea gggaaatggg tgetttatee 18660
aaatqqctcc aaqccaqqtq ggctactacc ttqttqttaq gggggtqtct tcctcacaac 18720
ctgtttttct cttcccagct gtggttggat caaggactca ttcctgcctt qqaqaaaata 18780
cttcaaccaq aqcaqqqaqc ctqqqqqtqt cqqqqcagqa qgctggqqat ggggqtggga 18840
tatgagggtg gcatgcagct gagggcaggg ccagggctgg tgtccctaag gttgtacaga 18900
ctcttgtgaa tatttgtatt ttccagatgg aataaaaagg cccgtgtaat taaccttcac 18960
catcagegee tagaateeeg gggggtaggg ggatggtata etttacagga tgacaatett 19020
gggagctaga actttgtagc cagagaaact tgggaggtct ggaatctcat gtgtctggag 19080
tcttggggaa gagaatctta gaagcagaaa accttggaac ataagaatct tggggagggt 19140
ctaggatett gaggagacca gateettgga catetaaaac ttgaaactag taggtetgea 19200
cccqaqaatt qcaqqqccaq tcatqcatac ccaaaqcctt caqcccatqq ccqaaattcc 19260
cttgctggac agggggcctt tcagcccctg cttggacgct tccagtaaca gggccctcac 19320
tqcaqqaatc qtqqqaqqqa qaqqqqcaqc acaqaqttqc tqqctqtcqq qqaaqqqaqq 19380
gagggccctg ggcagtccga gggccctgct gggcttgtgc ctcagggtgg gggctgcact 19440
cctccgcctt qcaqcctcct ggcctggtgc tgctgccagc cggaaggaca gtgacttcca 19500
gaggaaatge atattgatee tgettteage eteeggtggt ggetteteee aacceagete 19560
ttccctcctq agcctgcagc acggaggttt tggggggtcac tgctacctaa agaaggctaa 19620
ggccacttct gaggetggtc tgggagttta ctaaaggttc tgaagetggg ccgggctgcc 19680
cctgggatca ggagactcca gacagcagtc ctgacaatgg gaactacctc ctcagtcccc 19740
caaactggga ggtgtcccac agcagctgta ggattgtcct aggggtggag acctgagcac 19800
cttccactcc aaagcacagt atctgtgggc ctggcagtgg cctcagttcc cccatgagtg 19860
ccccggtccc ccaccccagg gtttccccac atcacatcca tccctgcttt gagaccccac 19920
tecceetgge etgttettta ttttgggtea etecettete ttteetggte atatetetee 19980
tgcaggccta coctgtgttg ggcccccag ccctgtctct gcatcgggtg cccccctgcc 20040
cetecttetg tecteageec ceteegeect tececetett gaggetgtaa tateegttte 20100
acgatttggg ggctgagttg ctataacaac agacggcgat tgtgttgtga agagcagctc 20160
getectgtge egectgeete etgtgetgee tecatecetg eageceagte ggtteetett 20220
qqctcctctc qtcactaccc tccaqttcca qtctqqcctc ttcctqqtqt qtqtqtqttqt 20280
gtgtgtgtgt gtgtgtat gcatgcatgc atatgtgtgt ccaggtctgc ctgcccggga 20340
tgtgacaagt agcggtcttc atggttgcat gtgtctgaat ttggtgtctg agcttcacat 20400
tgtatgcgcc tgtgtgcatg tgtgtgcatg gacatgcatg ctgtatctgc tgtgtttccc 20460
ctcccccatq tqtccccact qqcctttqca catqqqaqaa qqqcatqtqc tcaqcatatc 20520
acteaactgt ccacattggg tgggtacctg tgtgtggtgt gtgtgtgtgg ggggtgtgtc 20580
ttgaagtggc aggtcccaaa tgcttaggca atctgaacct tggaccttgc agagaggaga 20640
gatgteectg taggtgggag ggacagggag atgcagcage tgeecggtga cettttetge 20700
ccttgatggg caaagctggg ggtagggaaa ggagacaagt gctcatactt acctccctcc 20760
ctgcccaggc tectetgtaa gggtetgagt etgtetetgt gagecattge atetgtetgt 20820
ctatqccctq atqcctqqat qqacaaqqqq tqtqtqtqtq tqtqtqtqtq tqtqtqtqt 20880
agtgtgaggc tgcaggaaga ggaacagtgg gggatgggca ggaaagtggg ctgtggggtc 20940
                                                                 20951
agggaggcga t
```

<210> 4 <211> 609

<212> PRT

<213> Human

<400> 4

Met Ala Gly Thr Leu Asp Leu Asp Lys Gly Cys Thr Val Glu Glu Leu 1 5 10 15

Leu Arg Gly Cys Ile Glu Ala Phe Asp Asp Ser Gly Lys Val Arg Asp

			20					25					30		
Pro	Gln	Leu 35	Val	Arg	Met	Phe	Leu 40	Met	Met	His	Pro	Trp 45	Tyr	Ile	Pro
Ser	Ser 50	Gln	Leu	Ala	Ala	Lys 55	Leu	Leu	His	Ile	Tyr 60	Gln	Gln	Ser	Arg
Lys 65	Asp	Asn	Ser	Asn	Ser 70	Leu	Gln	Val	Lys	Thr 75	Cys	His	Leu	Val	Arg 80
-	_			85	Phe				90	_				95	
			100	_	Glu			105					110		
-	_	115			Leu		120		_			125		_	
_	130	_			Thr	135	_				140		_		
145					Phe 150	_				155					160
				165	Glu				170					175	
_	_		180		Val			185				_	190		
		195			Ser Lys		200					205			
	210				Val	215					220				
225					230 Val			_		235					240
				245	His				250					255	
_			260					265					270		
		275			Glu		280					285			
_	290		_		Ala Asp	295					300				
305					310					315					320
				325	Arg				330					335	
			340		Glu			345					350	_	
		355			Pro	_	360					365			
	370				Glu	375					380				_
385					Ser 390					395					400
		_		405	Val				410					415	
		-	420		Leu			425					430		
		435			Phe		440					445			
	450				Ile	455	_				460				
465	Азр	геп	ASP	GTIJ	Asn 470	GTII	нар	атА	cys	475	ser	Arg	GIU	GIU	Met 480

Val Ser Tyr Phe Leu Arg Ser Ser Ser Val Leu Gly Gly Arg Met Gly 490 485 Phe Val His Asn Phe Gln Glu Ser Asn Ser Leu Arg Pro Val Ala Cys 505 Arg His Cys Lys Ala Leu Ile Leu Gly Ile Tyr Lys Gln Gly Leu Lys 520 Cys Arg Ala Cys Gly Val Asn Cys His Lys Gln Cys Lys Asp Arg Leu 535 540 Ser Val Glu Cys Arg Arg Arg Ala Gln Ser Val Ser Leu Glu Gly Ser 550 555 Ala Pro Ser Pro Ser Pro Met His Ser His His Arg Ala Phe Ser 565 570 Phe Ser Leu Pro Arg Pro Gly Arg Arg Gly Ser Arg Pro Pro Glu Ile 585 Arg Glu Glu Val Gln Thr Val Glu Asp Gly Val Phe Asp Ile His Leu

<210> 5 <211> 664 <212> PRT

<213> Human

<400> 5 Gly Arg Gly Thr Gln Gly Trp Pro Gly Ser Ser Glu Gln His Val Gln 10 Glu Ala Thr Ser Ser Ala Gly Leu His Ser Gly Val Asp Glu Leu Gly Val Arg Ser Glu Pro Gly Gly Arg Leu Pro Glu Arg Ser Leu Gly Pro Ala His Pro Ala Pro Ala Ala Met Ala Gly Thr Leu Asp Leu Asp Lys 55 Gly Cys Thr Val Glu Glu Leu Leu Arg Gly Cys Ile Glu Ala Phe Asp Asp Ser Gly Lys Val Arg Asp Pro Gln Leu Val Arg Met Phe Leu Met 8.5 90 Met His Pro Trp Tyr Ile Pro Ser Ser Gln Leu Ala Ala Lys Leu Leu 105 His Ile Tyr Gln Gln Ser Arg Lys Asp Asn Ser Asn Ser Leu Gln Val 115 120 125 Lys Thr Cys His Leu Val Arg Tyr Trp Ile Ser Ala Phe Pro Ala Glu 135 140 Phe Asp Leu Asn Pro Glu Leu Ala Glu Gln Ile Lys Glu Leu Lys Ala 150 155 Leu Leu Asp Gln Glu Gly Asn Arg Arg His Ser Ser Leu Ile Asp Ile 170 Asp Ser Val Pro Thr Tyr Lys Trp Lys Arg Gln Val Thr Gln Arg Asn 185 Pro Val Gly Gln Lys Lys Arg Lys Met Ser Leu Leu Phe Asp His Leu 200 Glu Pro Met Glu Leu Ala Glu His Leu Thr Tyr Leu Glu Tyr Arg Ser 215 220 Phe Cys Lys Ile Leu Phe Gln Asp Tyr His Ser Phe Val Thr His Gly 230 235 Cys Thr Val Asp Asn Pro Val Leu Glu Arg Phe Ile Ser Leu Phe Asn

				245					250					255	
Ser	Val	Ser	Gln 260	Trp	Val	Gln	Leu	Met 265	Ile	Leu	Ser	Lys	Pro 270	Thr	Ala
Pro	Gln	Arg 275	Ala	Leu	Val	Ile	Thr 280	His	Phe	Val	His	Val 285	Ala	Glu	Lys
Leu	Leu 290	Gln	Leu	Gln	Asn	Phe 295	Asn	Thr	Leu	Met	Ala 300	Val	Val	Gly	Gly
Leu 305	Ser	His	Ser	Ser	Ile 310	Ser	Arg	Leu	Lys	Glu 315	Thr	His	Ser	His	Val 320
Ser	Pro	Glu	Thr	Ile 325	Lys	Leu	Trp	Glu	Gly 330	Leu	Thr	Glu	Leu	Val 335	Thr
Ala	Thr	Gly	Asn 340	Tyr	Gly	Asn	Tyr	Arg 345	Arg	Arg	Leu	Ala	Ala 350	Cys	Val
Gly	Phe	Arg 355	Phe	Pro	Ile	Leu	Gly 360	Val	His	Leu	Lys	Asp 365	Leu	Val	Ala
Leu	Gln 370	Leu	Ala	Leu	Pro	Asp 375		Leu	Asp	Pro	Ala 380	Arg	Thr	Arg	Leu
Asn 385	Gly	Ala	Lys	Met	Lys 390	Gln	Leu	Phe	Ser	Ile 395	Leu	Glu	Glu	Leu	Ala 400
	Val	Thr	Ser	Leu 405	Arg	Pro	Pro	Val	Gln 410		Asn	Pro	Asp	Leu 415	
Ser	Leu	Leu	Thr 420	Val	Ser	Leu	Asp	Gln 425	Tyr	Gln	Thr	Glu	Asp 430	Glu	Leu
Tyr	Gln	Leu 435	Ser	Leu	Gln	Arg	Glu 440	Pro	Arg	Ser	Lys	Ser 445	Ser	Pro	Thr
Ser	Pro 450	Thr	Ser	Cys	Thr	Pro 455	Pro	Pro	Arg	Pro	Pro 460	Val	Leu	Glu	Glu
Trp 465	Thr	Ser	Ala	Ala	Lys 470	Pro	Lys	Leu	Asp	Gln 475	Ala	Leu	Val	Val	Glu 480
His	Ile	Glu	Lys	Met 485	Val	Glu	Ser	Val	Phe 490	Arg	Asn	Phe	Asp	Val 495	Asp
Gly	Asp	Gly	His 500	Ile	Ser	Gln	Glu	Glu 505	Phe	Gln	Ile	Ile	Arg 510	Gly	Asn
Phe	Pro	Tyr 515	Leu	Ser	Ala	Phe	Gly 520	Asp	Leu	Asp	Gln	Asn 525	Gln	Asp	Gly
Cys	Ile 530	Ser	Arg	Glu	Glu	Met 535	Val	Ser	Tyr	Phe	Leu 540	Arg	Ser	Ser	Ser
545					550					555		Gln			560
Ser	Leu	Arg	Pro	Val 565	Ala	Суѕ	Arg	His	Cys 570	Lys	Ala	Leu	Ile	Leu 575	Gly
Ile	Tyr	Lys	Gln 580	Gly	Leu	Lys	Суѕ	Arg 585	Ala	Cys	Gly	Val	Asn 590	Cys	His
Lys	Gln	Cys 595	Lys	Asp	Arg	Leu	Ser 600	Val	Glu	Суѕ	Arg	Arg 605	Arg	Ala	Gln
Ser	Val 610	Ser	Leu	Glu	Gly	Ser 615	Ala	Pro	Ser	Pro	Ser 620	Pro	Met	His	Ser
His 625	His	His	Arg	Ala	Phe 630	Ser	Phe	Ser	Leu	Pro 635	Arg	Pro	Gly	Arg	Arg 640
Gly	Ser	Arg	Pro	Pro 645	Glu	Ile	Arg	Glu	Glu 650	Glu	Val	Gln	Thr	Val 655	Glu
Asp	Gly	Val	Phe 660	Asp	Ile	His	Leu								

<210> 6 <211> 608 <212> PRT <213> Mus musculus

Met Ala Ser Thr Leu Asp Leu Asp Lys Gly Cys Thr Val Glu Glu Leu Leu Arg Gly Cys Ile Glu Ala Phe Asp Asp Ser Gly Lys Val Arg Asp 25 Pro Gln Leu Val Arg Met Phe Leu Met Met His Pro Trp Tyr Ile Pro 40 Ser Ser Gln Leu Ala Ser Lys Leu Leu His Phe Tyr Gln Gln Ser Arg 55 Lys Asp Asn Ser Asn Ser Leu Gln Val Lys Thr Cys His Leu Val Arg Tyr Trp Val Ser Ala Phe Pro Ala Glu Phe Asp Leu Asn Pro Glu Leu 90 Ala Glu Pro Ile Lys Glu Leu Lys Ala Leu Leu Asp Gln Glu Gly Asn 105 Arg Arg His Ser Ser Leu Ile Asp Ile Glu Ser Val Pro Thr Tyr Lys 120 Trp Lys Arg Gln Val Thr Gln Arg Asn Pro Val Glu Gln Lys Lys Arg 135 Lys Met Ser Leu Leu Phe Asp His Leu Glu Pro Met Glu Leu Ala Glu 150 155 His Leu Thr Tyr Leu Glu Tyr Arg Ser Phe Cys Lys Ile Leu Phe Gln 170 Asp Tyr His Ser Phe Val Thr His Gly Cys Thr Val Asp Asn Pro Val 180 185 Leu Glu Arg Phe Ile Ser Leu Phe Asn Ser Val Ser Gln Trp Val Gln 200 Leu Met Ile Leu Ser Lys Pro Thr Ala Thr Gln Arg Ala Leu Val Ile 215 220 Thr His Phe Val His Val Ala Glu Lys Leu Gln Leu Gln Asn Phe 230 235 Asn Thr Leu Met Ala Val Val Gly Gly Leu Ser His Ser Ser Ile Ser 245 250 Arg Leu Lys Glu Thr His Ser His Val Ser Pro Asp Thr Ile Lys Leu 265 260 Trp Glu Gly Leu Thr Glu Leu Val Thr Ala Thr Gly Asn Tyr Ser Asn 280 Tyr Arg Arg Leu Ala Ala Cys Val Gly Phe Arg Phe Pro Ile Leu 295 300 Gly Val His Leu Lys Asp Leu Val Ala Leu Gln Leu Ala Leu Pro Asp 310 315 Trp Leu Asp Pro Gly Arg Thr Arg Leu Asn Gly Ala Lys Met Arg Gln 325 330 Leu Phe Ser Ile Leu Glu Glu Leu Ala Met Val Thr Ser Leu Arg Pro 340 345 Pro Val Gln Ala Asn Pro Asp Leu Leu Ser Leu Leu Thr Val Ser Leu 360 Asp Gln Tyr Gln Thr Glu Asp Glu Leu Tyr Gln Leu Ser Leu Gln Arg 375 Glu Pro Arg Ser Lys Ser Ser Pro Thr Ser Pro Thr Ser Cys Thr Pro 390 395 Pro Pro Arg Pro Pro Val Leu Glu Glu Trp Thr Ser Val Ala Lys Pro 405 410 Lys Leu Asp Gln Ala Leu Val Ala Glu His Ile Glu Lys Met Val Glu

```
425
           420
Ser Val Phe Arg Asn Phe Asp Val Asp Gly Asp Gly His Ile Ser Gln
                          440
Glu Glu Phe Gln Ile Ile Arg Gly Asn Phe Pro Tyr Leu Ser Ala Phe
           455
Gly Asp Leu Asp Gln Asn Gln Asp Gly Cys Ile Ser Arg Glu Glu Met
                 470
Ile Ser Tyr Phe Leu Arg Ser Ser Ser Val Leu Gly Gly Arg Met Gly
                    490
            485
Phe Val His Asn Phe Gln Glu Ser Asn Ser Leu Arg Pro Val Ala Cys
                             505
Arg His Cys Lys Ala Leu Ile Leu Gly Ile Tyr Lys Gln Gly Leu Lys
                                             525
                          520
       515
Cys Arg Ala Cys Gly Val Asn Cys His Lys Gln Cys Lys Asp Arg Leu
                                          540
                       535
Ser Val Glu Cys Arg Arg Arg Ala Gln Ser Val Ser Leu Glu Gly Ser
                                      555
                  550
Ala Pro Ser Pro Ser Pro Thr His Thr His His Arg Ala Phe Ser Phe
                                  570
              565
Ser Leu Pro Arg Pro Gly Arg Arg Ser Ser Arg Pro Pro Glu Ile Arg
                              585
Glu Glu Glu Val Gln Thr Val Glu Asp Gly Val Phe Asp Ile His Leu
                           600
<210> 7
<211> 591
<212> PRT
<213> Human
<400> 7
Gly Ser Ser Gly Leu Gly Lys Ala Ala Thr Leu Asp Glu Leu Leu Cys
                                  10
```

Thr Cys Ile Glu Met Phe Asp Asp Asn Gly Glu Leu Asp Asn Ser Tyr 25 Leu Pro Arg Ile Val Leu Leu Met His Arg Trp Tyr Leu Ser Ser Thr 40 Glu Leu Ala Glu Lys Leu Leu Cys Met Tyr Arg Asn Ala Thr Gly Glu 60 55 Ser Cys Asn Glu Phe Arg Leu Lys Ile Cys Tyr Phe Met Arg Tyr Trp 70 75 Ile Leu Lys Phe Pro Ala Glu Phe Asn Leu Asp Leu Gly Leu Ile Arg 90 Met Thr Glu Glu Phe Arg Glu Val Ala Ser Gln Leu Gly Tyr Glu Lys 105 100 His Val Ser Leu Ile Asp Ile Ser Ser Ile Pro Ser Tyr Asp Trp Met 120 Arg Arg Val Thr Gln Arg Lys Lys Val Ser Lys Lys Gly Lys Ala Cys 135 Leu Leu Phe Asp His Leu Glu Pro Ile Glu Leu Ala Glu His Leu Thr 155 150 Phe Leu Glu His Lys Ser Phe Arg Arg Ile Ser Phe Thr Asp Tyr Gln 170 165 Ser Tyr Val Ile His Gly Cys Leu Glu Asn Asn Pro Thr Leu Glu Arg 185 Ser Ile Ala Leu Phe Asn Gly Ile Ser Lys Trp Val Gln Leu Met Val 200

```
Leu Ser Lys Pro Thr Pro Gln Gln Arg Ala Glu Val Ile Thr Lys Phe
                       215
                                           220
Ile Asn Val Ala Lys Lys Leu Leu Gln Leu Lys Asn Phe Asn Thr Leu
                   230
                                      235
Met Ala Val Val Gly Gly Leu Ser His Ser Ser Ile Ser Arg Leu Lys
               245
                                  250
Glu Thr His Ser His Leu Ser Ser Glu Val Thr Lys Asn Trp Asn Glu
                               265
           260
Met Thr Glu Leu Val Ser Ser Asn Gly Asn Tyr Cys Asn Tyr Arg Lys
                           280
Ala Phe Ala Asp Cys Asp Gly Phe Lys Ile Pro Ile Leu Gly Val His
                       295
Leu Lys Asp Leu Ile Ala Val His Val Ile Phe Pro Asp Trp Thr Glu
                   310
Glu Asn Lys Val Asn Ile Val Lys Met His Gln Leu Ser Val Thr Leu
                                   330
               325
Ser Glu Leu Val Ser Leu Gln Asn Ala Ser His His Leu Glu Pro Asn
                              345
Met Asp Leu Ile Asn Leu Leu Thr Leu Ser Leu Asp Leu Tyr His Thr
                           360
Glu Asp Asp Ile Tyr Lys Leu Ser Leu Val Leu Glu Pro Arg Asn Ser
                      375
                                          380
Lys Ser Pro Thr Ser Pro Thr Thr Pro Asn Lys Pro Val Val Pro Leu
                  390
                                      395
Glu Trp Ala Leu Gly Val Met Pro Lys Pro Asp Pro Thr Val Ile Asn
               405
                                  410
Lys His Ile Arg Lys Leu Val Glu Ser Val Phe Arg Asn Tyr Asp His
                               425
                                                  430
Asp His Asp Gly Tyr Ile Ser Gln Glu Asp Phe Glu Ser Ile Ala Ala
                           440
Asn Phe Pro Phe Leu Asp Ser Phe Cys Val Leu Asp Lys Asp Gln Asp
                       455
                                           460
Gly Leu Ile Ser Lys Asp Glu Met Met Ala Tyr Phe Leu Arg Ala Lys
                   470
                                       475
Ser Gln Leu His Cys Lys Met Gly Pro Gly Phe Ile His Asn Phe Gln
               485
                                   490
Glu Met Thr Tyr Leu Lys Pro Thr Phe Cys Glu His Cys Ala Gly Phe
                               505
           500
Leu Trp Gly Ile Ile Lys Gln Gly Tyr Lys Cys Lys Asp Cys Gly Ala
                           520
Asn Cys His Lys Gln Cys Lys Asp Leu Leu Val Leu Ala Cys Arg Arg
                      535
                                          540
Phe Ala Arg Ala Pro Ser Leu Ser Ser Gly His Gly Ser Leu Pro Gly
                   550
                                       555
Ser Pro Ser Leu Pro Pro Ala Gln Asp Glu Val Phe Glu Phe Pro Gly
                                   570
               565
Val Thr Ala Gly His Arg Asp Leu Asp Ser Arg Ala Ile Thr Leu
                               585
```

```
<210> 8
```

<sup>&</sup>lt;211> 581

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Rattus norvegicus

<sup>&</sup>lt;400> 8

Gly Ser Arg Ala Gly Pro Lys Gly Arg Leu Glu Ala Lys Ser Thr Asn

1				5					10					15	
Ser			Pro 20	Ala				25	Ala				30		
		35	Ser				40					45			
	50		Ser			55					60				
65	Ser		Gln		70					75					80
			Ala	85					90					95	
			Lys 100					T02					110		
		115	Trp				120					125			
	130		Ser			135					140				
145			Ser		150					155					T00
			Ser	165					170					1/5	
			Lys 180					185					190		
		195	His				200					205			
	21.0		Asp Met			215					220				
225					230					235					240
			Leu	245					250					255	
			Ile 260					265					270		
		275	Asn				280					285			
	290		Arg			295					300				
305					310					315					Asn 320
				325	,				330	)				333	
			340	)				345	5				350	1	Ala
		355	5				360	)				365	<b>)</b>		Leu
	370	)				375	5				380	)			Ala
385	5				390	)				395					Ser 400
				405	5				410	)				415	
			420	)				425	5				430	)	Pro
		43	5				44(	)				445	5		Pro
Lys	5 Thi 450		e Se:	r Lys	s His	s Val 45	L Glr 5	n Ar	g Me	t Va.	L Asp 460	) Sei	r val	r rne	e Lys

#### CL001165

Asn Tyr Asp Leu Asp Gln Asp Gly Tyr Ile Ser Gln Glu Glu Phe Glu 470 475 Lys Ile Ala Ala Ser Phe Pro Phe Ser Phe Cys Val Met Asp Lys Asp 485 490 Arg Glu Gly Leu Ile Ser Arg Asp Glu Ile Thr Ala Tyr Phe Met Arg 505 510 500 Ala Ser Ser Ile Tyr Ser Lys Leu Gly Leu Gly Phe Pro His Asn Phe 515 520 525 Gln Glu Thr Thr Tyr Leu Lys Pro Thr Phe Cys Asp Asn Cys Ala Gly 535 Phe Leu Trp Gly Val Ile Lys Gln Gly Tyr Arg Cys Lys Asp Cys Gly 550 555 545 Met Asn Cys His Lys Gln Cys Lys Asp Leu Val Val Phe Glu Cys Lys 565 570 Lys Arg Ser Lys Ser 580